

Recent Studies on a μ -Opioid Receptor Purified from Bovine Striatum^a

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The existence of opioid receptors in the central nervous system of animals and humans was demonstrated in 1973 simultaneously by our laboratory^{1,2} and by others.³⁻⁵ At the time only a single type of receptor was postulated, a concept soon challenged and disproved. Martin and co-workers^{6,7} postulated three classes of receptors, μ , κ , and σ . The latter is now not considered to be an opioid receptor and has been replaced by δ , an enkephalin-preferring receptor first demonstrated by Kosterlitz and co-workers⁸ in the mouse *vas deferens*. The enkephalins are therefore thought to be the endogenous ligands of the δ -receptor, dynorphin that of the κ -receptor, and β -endorphin a putative endogenous ligand of the μ -receptor. Within the past year the δ -opioid receptor was cloned independently by two laboratories.^{9,10} This was rapidly followed by the cloning of a μ - and a κ -receptor,^{11,12} by low stringency screening of cDNA libraries with oligonucleotide probes from the structure of the δ -receptor cDNA. The protein sequences derived from the cDNA's coding for the three types of opioid receptors are shown in FIGURE 1. All of them have structures with seven putative transmembrane domains, typical of G-protein coupled receptors. The figure shows a number of receptors of this class that exhibit high levels of homology with the opioid receptors.

There is considerable pharmacological evidence for the existence of multiple subtypes of each of the three major types of opioid receptors. However, no evidence for subtypes has yet been provided by cloning, that is, all clones of a given type, obtained in a number of laboratories,¹³⁻¹⁹ including those cited earlier, have proved to be identical to the cDNA's coding for the μ -, δ -, and κ -opioid receptors originally published. The present paper discusses recent work in our laboratory on a μ -opioid receptor purified to homogeneity from bovine striatal membranes, which appears to be a subtype, different from the μ -receptor which has been cloned in several laboratories. Final proof of this hypothesis awaits the cloning of this protein.

RESULTS AND DISCUSSION

Purification and Microsequencing of a μ -Opioid Binding Protein

The purification to homogeneity of an opioid binding protein (OBP) from bovine striatal membrane was reported by us some years ago.²⁰ Briefly, the membranes were solubilized with digitonin and the extract was purified in two steps, ligand-affinity chromatography, which provided the major purification (4000- to 5000-fold) fol-

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lowed by lectin-affinity chromatography on wheat-germ agglutinin agarose, for a total purification of 60,000- to 70,000-fold. The purified protein gives a single band on SDS-polyacrylamide gel electrophoresis corresponding to a molecular mass of 65 kDa. It is highly glycosylated and enzymatic deglycosylation yields a protein of 40–43 kDa (Gioannini and Simon, unpublished results). Preliminary evidence suggested that it has the properties of a μ -opioid receptor, though this evidence was not conclusive because the purified protein bound only antagonists with high affinity. High-affinity agonist binding was lost, presumably due to uncoupling of the receptor protein from G-protein (see below for experiments on reconstitution).

The purified protein was found to be blocked at the N-terminus. Fragmentation with cyanogen bromide and sequencing, kindly carried out for us by Dr. C. Strader at Merck & Co., provided two peptides, one 20 amino acids long (peptide I) and the other 13 amino acids in length (peptide II). More recently, Dr. W. Burgess at the American Red Cross obtained four more short peptides for us by fragmentation with lysine-protease and microsequencing. It is of interest that none of these six peptides are found in the published amino acid sequence of the cloned μ -opioid receptor. Attempts to clone OBP, using PCR and direct screening of cDNA and genomic libraries, with oligonucleotides made from peptides I and II, have not yet succeeded. This is presumably because of the relatively short peptides and their high level of degeneracy. The following sections deal with studies using antibodies (Abs) made against portions of peptide I and II and with the reconstitution of purified OBP with G-proteins in liposomes.

Antisera against Peptides Derived from Opioid Binding Protein

Our laboratory, in collaboration with Dr. Lawrence Taylor in Dr. Huda Akil's laboratory, University of Michigan (Ann Arbor, MI), produced six antisera against three sequences in peptides I and II.²¹ TABLE 1 lists the sequences and the Abs made against them. All of these Abs recognized the purified OBP, from which the peptides are derived. They immunoprecipitate the bulk of radioiodine-labeled OBP and give an immunoblot at the correct molecular mass of 65 kDa. Most of our subsequent studies were done with antiserum Ab165, made against the N-terminal 12 amino acids of peptide I, the most immunoreactive serum against the OBP protein. In studies of a number of bovine brain regions, positive immunoblot signals corresponding to a molecular mass of 65 kDa were obtained with regions known to be rich in μ -receptors, whereas no signal was obtained with regions devoid of opioid receptors, such as white matter. Similarly, the cell line, SK-N-SH, high in μ -opioid receptors, gave a band in immunoblots, whereas HeLa and C-6 cells, devoid of opioid receptors, do not. Interestingly, NG-108-15 cells that express only δ -receptors give a band at a slightly lower molecular mass of approximately 60 kDa with Ab165. This may suggest cross-reaction with δ -receptors or the presence of silent and slightly different μ -receptors. Experiments are in progress to determine which of these alternatives is correct.

More recently, Dr. Hiller in our laboratory has carried out immunocytochemical studies of the distribution of Ab165-reactive material in the rat central nervous system.²² It parallels very closely the distribution of μ -opioid receptor binding, determined by autoradiography by ourselves²³ and by others.^{24,25} FIGURE 2 shows an example of these studies, namely, the spinal cord distribution of immunoreactivity to Ab165.

	1	#	#	#	50
MUOR1MDS	STGPGNTSDC	SDPLAQASCS	PAPGSWLNL	HVDGNQSDPC
DOR1	MELVPSARAE	LQSSPLVNLSDAFPS
KOR1MESP	IQIFRGDGP	TCSPSACLLP	NSSSWFPNWA
SOMR1MFPNAPP	LPHSSPSSSP	GGCGEGVCSR
FPEP
OBPRMASP	AGNLS..AWPG	WGWP..PAA	LRNLTSSPAP
NEUROKR	MASVPRGENW	TDGTVEVGTH	TGNLSSALGV	TEWLALQAGN	FSSALGLPAT
B2ARMEP
	51#	#			100
MUOR1	GLNRTGLGGN	DSLCPQTG..	...SPSMVTA	ITIMALYIY	CYVGLGNYL
DOR1	AFPSAGANAS	GSPGARS..	...S..SLALA	IAITALYSAV	CAVGLGNYL
KOR1	ESDSNGSVGS	EDQQLESA..	...HISPAIP	VIITAVYSV	FVVGVLGNSL
SOMR1	GPSSGAADGM	EEPGRNSSQN	GTLSEGGQSA	ILISFIYSV	CLVGLCGNSL
FPEP	...METNSS	LPTNISGGTP	AVSAGYLFLD	IITYLVFAVT	FVLQVLGNSL
OBPR	TASPSAPSW	TPSPRPGPAH	PFLQPPWAVA	LWSLA..YGAV	VAVAVLGNLV
NEUROKR	TQAPSQV...	...RANLTN	QFVQPSWRIA	LWSLA..YGLV	VAVAVFGNLI
B2AR	HGNDSDFLLA	PNGSRAPGHD	ITQERDEAWV	VGMALMSVI	VLAIVFGNVL
	101				150
MUOR1	VMYVIVRYTK	MKTATNIYIF	NLALADALAT	STLP..FQSAK	YLMGTWPFGT
DOR1	VMFGIVRYTK	LKTATNIYIF	NLALADALAT	STLP..FQSAK	YLMGTWPFGE
KOR1	VMFVIIRYTK	MKTATNIYIF	NLALADALVT	TTMP..FQSAV	YLMNSWPFGE
SOMR1	VIVVILRYAK	MKTATNIYIL	NLAIADDELM	LSVP..FLVTS	TLLRHWPFGA
FPEP	VIWVAG..FRM	THVTITISYL	NLAVADFCFT	STLPFFHVRK	AMGGHWPFGW
OBPR	VIWIVLAHKK	MRTVTNSFLV	NLAFADAAMA	ALNALVNFY	ALHGEWYFGA
NEUROKR	VIWIVLAHKK	MRTVTNYFLV	NLAFSDASVA	AFNTLINFIY	GLHSEWYFGA
B2AR	VITAIKFER	LQTVTNFIT	SLACADLVHG	LAVVPPGASH	ILMKMNWFCN
	151				200
MUOR1	ILCKIVISID	YNNMFTSIFT	LTMMSVDRI	AVCHPVKALD	FRTFRNAKIV
DOR1	LLCKAVLSID	YNNMFTSIFT	LTMMSVDRI	AVCHPVKALD	FRTPAKAKII
KOR1	VLCXIVISID	YNNMFTSIFT	LTMMSVDRI	AVCHPVKALD	FRTPLKAKII
SOMR1	LLCRLVLSVD	AVNMFTSIYC	LTVLSVDRI	AVEHPIKAAR	YRRPTVAKVV
FPEP	FLCKFVFTIV	DINLFGSVFL	IALIALDRCV	CVLHPVWTON	HRTVSLAKKV
OBPR	NYCRFQNFPP	ITAVFASIYS	MTAIAVDRI	AIIDPLKPR	..LSATATRIV
NEUROKR	NYCRFQNFPP	ITAVFASIYS	MTAIAVDRI	AIIDPLKPR	..LSATATKIV
B2AR	FWCEFTSID	VLCVTASIE	LCVIAVDRI	AITSPFKYQS	LLTKNKARV
	201				250
MUOR1	MVCNVLSSA	YGLPVPMAT	TKYRQ..GSI	DCTL..TFS..	..HPTW..YWE
DOR1	NICIWVLASG	VGVPIMVMV	TQPRD..GAV	VCML..QFP..	..SPSW..YWD
KOR1	NICIWLLASS	VGISAIVLGG	TKVREDVDVI	ECSL..QFP..	..DDEYSWD
SOMR1	NLGVVLSLL	VILPIVVFSS	TAANS..GTV	ACNM..LMP..	..EPAQRWLV
FPEP	IICPVVMALL	LTLPIVIRVT	T..VPGKTGT	ACTF..NFSPW	TNDPKERIKV
OBPR	IGSIWILAF	LAFPQCCLYSK	IKVMP.....	GRTL..CYVQW	PEGSRQHFTY
NEUROKR	IGSIWILAF	LAFPQCCLYSK	IKVMP.....	GRTL..CYVQW	PEGPKQHFTY
B2AR	ILMVIVSGL	TSFLPIQMHV	YRATHKQAI	CYAKETCCDF	FTNQAYAIAS

FIGURE 1. Amino acid sequences of the cloned μ -, δ -, and κ -opioid receptors and homologous G-protein coupled receptors. Boldface type and shading: putative transmembrane domains. *, putative sites for phosphorylation by protein kinase A. #, putative sites for N-linked glycosylation. (Adapted from Wang *et al.*,¹³ with permission.)

Reconstitution of High-Affinity Binding and GTPase Stimulation

Our preliminary pharmacological evidence as well as the results of the studies with the antibodies directed against peptides from purified OBP suggested that OBP was a μ -binding protein. However, these results were not conclusive, and real proof had to come from pharmacological studies of the reconstituted purified OBP, described in this section.

	251				300
MUOR1	NLLKICV...	..FIFAFVVF	ILIIIVCYGL	MILRLKSVRM	LSGS....KE
DOR1	TVTKICV...	..FLFAPVVF	ILIIIVCYGL	MILRLRSVRL	LSGS....KE
KOR1	LFMKICV...	..FVFAPVIP	VLIIVCYTL	MILRLKSVRL	LSGS....RE
SOMR1	GFV.LYT...	..FLMGFLLP	VGAICLCYVL	IIAKHRMVPS	RPAG.....S
FPEP	AVAMLTVRGI	IRFIIGFSAP	MSIVAVSYGL	IATKIHKQGL	IKSS.....
OBPR	HMIIV.....	..VLVYCFP	LLIMGITYTI	VGITLWCGEI	PGDTCCKYQE
NEUROKR	HIIIV.....	..ILVYCFP	LLIMGVITYI	VGITLWCGEI	PGDTCCKYHE
B2AR	SIVSFYVP..	..LVVMVFVYS	RVFQVAKRQL	QKIDKSEGRF	HAQNLSQVEQ
	301				350
MUOR1	KDRN.....LRR	ITRMVLVVVA	VFIICWPIH	IFVIVWTLVD
DOR1	KDRS.....LRR	ITRMVLVVVG	AFVVCWAPIH	IFVIVWTLVD
KOR1	KDRN.....LRR	ITKLVLVVVA	VFIICWPIH	IFILVEALGS
SOMR1	TQRS.....ERK	ITLMVMVMVM	VFVICWMPFY	VVQLVNVFAE
FPEPR	PLRVLSFVAA	AFFLCWSPIQ	VVALIATVRI
OBPR	QLKA.....KRK	VVKMMIIVVV	TFAICWLPYH	IYFILTAIYQ
NEUROKR	QLKA.....KRK	VVKMMIIVVV	TFAICWLPYH	VYFILTAIYQ
B2AR	DGRSGHGLRS	SSKFCLEKHK	ALKTLGIIMG	TPTLCWLPFF	IVNIVHVIRA
	351				400
MUOR1	I.PETTF...	QTVSWHFCIA	LGYNSSCLNP	VLYAFDENF	KRCF.REF..
DOR1	INRRDPL...	VVAALHLCIA	LGYNSSCLNP	VLYAFDENF	KRCF.RQL..
KOR1	TSHSTA...	ALSSYYFCIA	LGYNSSCLNP	VLYAFDENF	KRCF.RDF..
SOMR1	QDDATVS...	Q.....LSVI	LGYNSSCLNP	ILYGFSLDNF	KRSFQRIIL..
FPEP	RELLQGMKE	IGIAVDVTS	LAFFNSCLNP	MLYVFMQDF	RERLIHAL..
OBPR	QLNRWKYIQ	VYLASFV...	LAMSSTHYNP	IYCCCLNKR	RAGFKRAFWR
NEUROKR	QLNRWKYIQ	VYLASFV...	LAMSSTHYNP	IYCCCLNKR	RAGFKRAFWR
B2AR	NLIPKEV...YILLNW	LCYVNSAFNP	LIYC.RSPDF	RIAFQELLCL
	401				450
MUOR1	.CIPTSSTIE	QONSTRVRQN	TREHPSTANT	VDRTNHQLN	LEARTAPLP.
DOR1	.CRTPCGRQE	PGSLRRPRQA	TTRERVACT	PS.....DG	PGGGAAA...
KOR1	.CFPIKMRME	RQSTNRVR.N	TVQDPASMRD	VGGMKNFV..
SOMR1	.CL...SWMD	NAAEPPVDYY	ATALKSRAYS	VEDFQENLE	SGGVFRNGTC
FPEP	.PASLERALT	EDSTQTSDTA	TNSTLPSAEV	ALQAK.....
OBPR	CPFIHVSSYD	ELELKATRLH	PMRQ.SSLYT	VTRMESMSV	FDSNDGDSAR
NEUROKR	CPFIQVSSYD	ELELKATTRFH	PTRQ.SSLYT	VSRMESVTVL	FDPNDGDPTK
B2AR	RRSSSKTYGN	GYSSNSNGRT	DYTGEOQAYQ	LCQEKENELL	CEEAPGMEGF

FIGURE 1. (continued)

Our initial studies of OBPR reconstitution²⁶ were done with a CHAPS extract of bovine striatal membranes as the source of phospholipids and G-proteins. This extract was prepared in the absence of sodium, which we have shown to solubilize very little opioid binding activity. Heating the extract at 37 °C for 30 min removed any residual traces of opioid binding activity. Lipid vesicles were formed by adding purified OBPR to the CHAPS extract, precipitating the mixture with polyethyleneglycol-6000 (PEG), and resuspending the pellet, resulting from centrifugation at 12,000 × g.

TABLE 1. Antipeptide Antibodies Generated against Amino Acid Sequences Derived from Opioid Binding Protein

	Amino Acid Sequence	Antibody
Peptide I-1-12	IRNLRQDRSKYY[X]	165, 166, 6639
Peptide I-14-20	NFFYKRL	163
Peptide II-1-9	Y*SNNVLFVSH[XFND]	161, 162

NOTE: *, Y-tyrosine added to permit iodination of peptide. X indicates unknown amino acids. Bracketed residues were not used for antibody production. (Gioannini *et al.*²¹ Reprinted with permission from *Molecular Pharmacology*.)

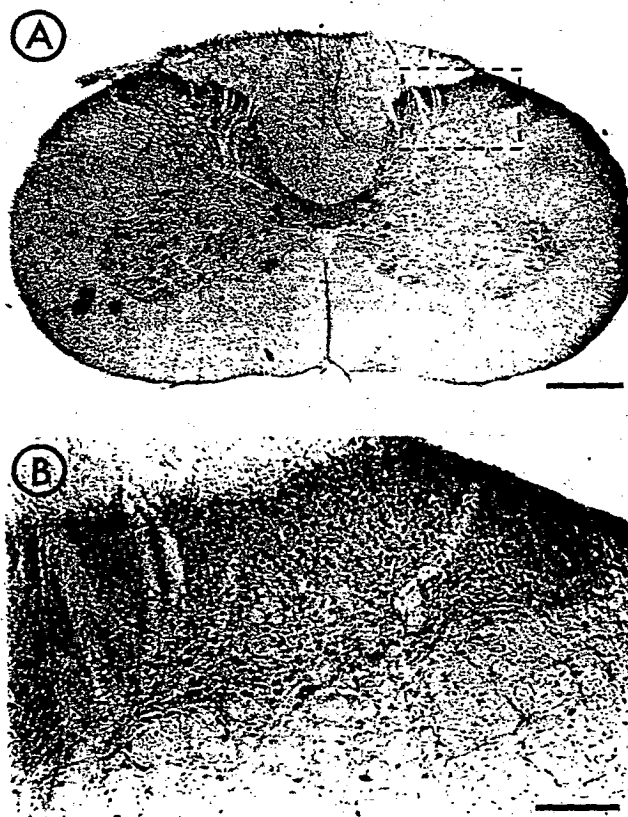
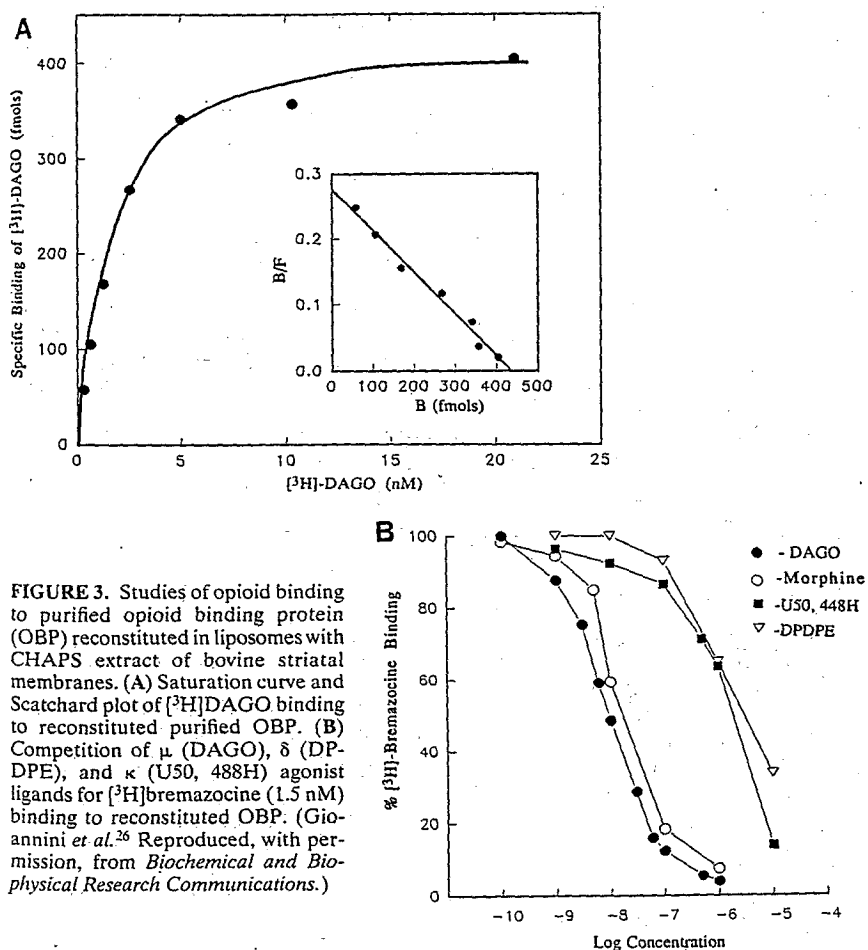


FIGURE 2. Distribution of immunoreactivity against Ab165 in the rat lumbar spinal cord. (A) High levels of immunoreactivity are seen in lamina I and lamina II (substantia gelatinosa) of the dorsal horn. Higher magnification of the (boxed area of panel A) dorsal horn, shown in panel B, reveals that laminae III and IV also contain immunopositive cell bodies. An edge artifact is apparent in the upper right corner of this panel. It is also notable that the median aspect of lamina V (see panel A) contains a substantial level of immunoreactivity. In the ventral horns, large individual immunopositive cell bodies can be seen in laminae VII, VIII, and IX. Bars, A = 100 μ m, B = 2 μ m. (Hiller *et al.*²² Reproduced, with permission, from *Neuroscience*.)

in Tris buffer containing magnesium. FIGURE 3 shows that the selective μ -agonist DAGO binds saturably and with high affinity to the reconstituted OBP. The figure also shows that the binding is very selective for μ -ligands. The binding was found to be completely dependent on and proportional to the amount of OBP added to the liposomes, stereospecific and sensitive to inhibition by GTP γ S. The affinity of DAGO for reconstituted OBP was 1.5 nM, identical to that seen in the membrane-bound μ -opioid receptor in bovine striatal membranes. Functional coupling was also achieved, as evidenced by the stimulation of low K_m GTPase by μ -agonists, such as DAGO.

More recently we repeated the reconstitution studies with purified G-proteins in collaboration with the laboratory of John Hildebrandt at South Carolina Medical University (Charleston, SC) (manuscript in preparation). Most of the studies were carried out with a highly purified mixture of brain G-proteins, composed of 80% G_{oA} , 7-8% G_{oC} , 7-8% G_{i1} and small amounts of G_{oB} and G_{i2} . Some experiments with individual purified G-proteins were also done. The liposomes were formed by mixing purified OBP with phosphatidylcholine (Sigma) and the purified G-proteins. The mixture was precipitated with PEG, centrifuged at $100,000 \times g$, and resuspended in Tris buffer containing magnesium. Functional reconstitution was assessed by measurement of opioid stimulation of low K_m GTPase. Opioid ligands, such as DAGO, morphine and levorphanol, produced up to 100% stimulation of GTPase. The stimulation was reversed by (-)naloxone but not by (+)naloxone. Dextrorphan, the inactive enantiomer of levorphanol, was inactive and so were δ - and κ -agonists.



Studies of μ -agonist binding showed that high-affinity binding to reconstituted OBP was obtained both with the G-protein mixture and with individual G-proteins. Differences in efficacy between different G-proteins were not significant. Furthermore, differences observed could represent differences in incorporation of various G-proteins into liposomes, in the level of coupling or in the nature of the coupling. We cannot at present distinguish among these possibilities. It should be noted that the level of binding and the affinity of DAGO binding were lower than those seen when CHAPS extract was the source of G-proteins. This could be due to the absence of factors that favor high levels of incorporation into liposomes and coupling or to some slight denaturation of G-proteins during purification. The affinity of DAGO binding, $K_d = 7$ nM, was high but somewhat lower than the $K_d = 1.5$ nM we observed in the experiments with CHAPS extract.

In summary, our studies with antisera directed against peptides derived from purified OBP, and especially the pharmacological profile obtained with reconstituted OBP, demonstrate clearly that we have purified a μ -receptor. The fact that none of the peptides obtained from the purified protein are identical to sequences in the cloned μ -receptor suggests that it may be a different subtype, the predominant subtype in bovine brain. Our results suggest that this subtype of μ -opioid receptor is also present in rat brain or, alternatively, that sequence homology and/or tertiary structural similarities give rise to cross-reactivity of the rat μ -receptor with our antisera.

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